

XP-000144843

Q9y4b8.Trembl

P.D. 1-11-1999

P. 1



!!AA_SEQUENCE 1.0

ID Q9Y4B8 PRELIMINARY; PRT; 468 AA.
AC Q9Y4B8;
DT 01-NOV-1999 (TrEMBLrel. 12, Created)
DT 01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
DT 01-MAY-2000 (TrEMBLrel. 13, Last annotation update)
DE KIAA0772 PROTEIN.
GN KIAA0772.
OS Homo sapiens (Human).
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
RN [1]
RP SEQUENCE FROM N.A.
RC TISSUE=BRAIN;

RX MEDLINE; 99087487.
RA Nagase T., Ishikawa K., Suyama M., Kikuno R., Miyajima N., Tanaka A.,
RA Kotani H., Nomura N., Ohara O.;
RT "Prediction of the coding sequences of unidentified human genes. XI.
RT The complete sequences of 100 new cDNA clones from brain which code
RT for large proteins in vitro.";
RL DNA Res. 5:277-286(1998).
OR EMBL; AB018315; BAA34492.1; -.
DR INTERPRO; IPR000648; -.
DR PFAM; PF01237; Oxysterol_BP; 1.
DR PROSITE; PS01013; OSBP; 1.
SQ SEQUENCE 468 AA; 53971 MW; 9C51392E0EDF4DF8 CRC64;

Q9y4b8 Length: 468 August 11, 19100 11:16 Type: P Check: 5128 ..

1 MNGEEFFDA VTEANQKVTG MIDLDTSKNN RIGKTGERPS QENGIQKHRT

51 SLPAPMFSRS DFSVWTILKK CVGLELSKIT MPIAFNEPLS FLQRITEYME
101 HVYLIHRASC QPQPLERMQS VAAFAVSAVA SQWERTGKPF NPLLGETYEL
151 IREDLGFRFI SEQVSHHPPI SAFHSEGLNH DFLFHGSIYP KLKFWGKSVE
201 AEPRGTITLE LLKHNEAYTW TNPTCCVHNV IIGKLWIEQY GTVEILNHRT
251 GHKCVLHFKP CGLFGKELHK VEGHIQDKNK KKLFIYWKW TECLWGIDPV
301 SYESFKKQER RGDHLRKAKL DEDSGKADSD VADDVPVAQE TVQVIPGSKL
351 LWRINTRPPN SAQMYNFTSF TVSLNELETG MEKTLPPPTDC RLRPDIRGME
401 NGNMDLASQE KERLEEKQRE ARRERAKEEA EWQTRWFYPG NNPYTGTPDW
451 LYAGDYFERN FSDCPDIY